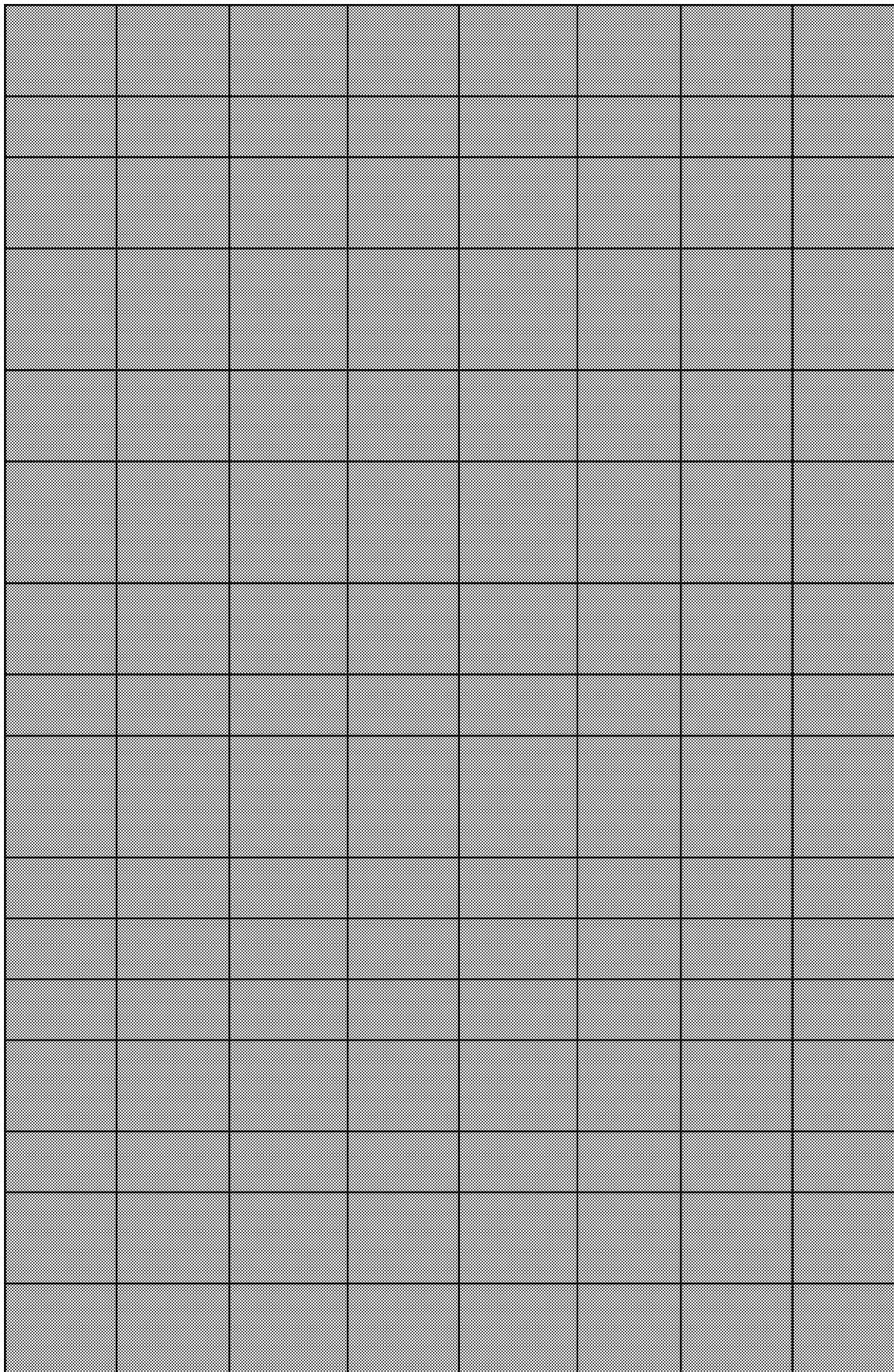


Level 1



2309
2310
2311
2312
2313
2314
2315
2316
2317
2318
2319
2320
2321
2322
2323
2324
2325
2326

S. Lopez-Gomollon, E. Sevilla, M. T. Bes, M. L. Peleato, M. F. Fillat. New insights into the role of Fur proteins: FurB (AlI2473) from Anabaena protects DNA and increases cell survival under oxidative stress. Biochem J. 2009. 418:201-7
L. Z. Zhou, A. P. Johnson, T. A. Rando. NF kappa B and AP-1 mediate transcriptional responses to oxidative stress in skeletal muscle cells. Free Radic Biol Med. 2001. 31:1405-16
B. W. Yun, M. J. Skelly, M. Yin, M. Yu, B. G. Mun, S. U. Lee, A. Hussain, S. H. Spoel, G. J. Loake. Nitric oxide and S-nitrosoglutathione function additively during plant immunity. New Phytol. 2016. 211:516-26
S. I. Liochev, A. Hausladen, I. Fridovich. Nitroreductase A is regulated as a member of the soxRS regulon of Escherichia coli. Proc Natl Acad Sci U S A. 1999. 96:3537-9
A. Muller, J. H. Hoffmann, H. E. Meyer, F. Narberhaus, U. Jakob, L. I. Leichert. Nonnative disulfide bond formation activates the sigma32-dependent heat shock response in Escherichia coli. J Bacteriol. 2013. 195:2807-16
J. Prosecka, A. V. Orlov, Y. S. Fantin, V. V. Zinchenko, M. M. Babykin, M. Tichy. A novel ATP-binding cassette transporter is responsible for resistance to viologen herbicides in the cyanobacterium Synechocystis sp. PCC 6803. FEBS j. 2009. 276:4001-11
J. H. Niazi, B. C. Kim, J. M. Ahn, M. B. Gu. A novel bioluminescent bacterial biosensor using the highly specific oxidative stress-inducible pgi gene. Biosens Bioelectron. 2008. 24:670-5
E. S. Seong, M. H. Wang. A novel CaAbs1 gene induced by early-abiotic stresses in pepper. BMB Rep. 2008. 41:86-91
S. Tsukamoto, S. Morita, E. Hirano, H. Yokoi, T. Masumura, K. Tanaka. A novel cis-element that is responsive to oxidative stress regulates three antioxidant defense genes in rice. Plant Physiol. 2005. 137:317-27
C. Li, J. Tao, D. Mao, C. He. A novel manganese efflux system, YebN, is required for virulence by <i>Xanthomonas oryzae</i> pv. <i>oryzae</i> . PLoS One. 2011. 6:e21983
D. Holland, Z. Faltin, A. Perl, G. Ben-Hayyim, Y. Eshdat. A novel plant glutathione peroxidase-like protein provides tolerance to oxygen radicals generated by paraquat in Escherichia coli. FEBS Lett. 1994. 337:52-5
D. G. Kehres, M. L. Zaharik, B. B. Finlay, M. E. Maguire. The NRAMP proteins of <i>Salmonella typhimurium</i> and <i>Escherichia coli</i> are selective manganese transporters involved in the response to reactive oxygen. Mol Microbiol. 2000. 36:1085-100
A. M. Pickering, R. A. Linder, H. Zhang, H. J. Forman, K. J. Davies. Nrf2-dependent induction of proteasome and Pa28alpha/beta regulator are required for adaptation to oxidative stress. J Biol Chem. 2012. 287:10021-31
M. I. Ramirez-Diaz, A. Diaz-Magana, V. Meza-Carmen, L. Johnstone, C. Cervantes, C. Rensing. Nucleotide sequence of <i>Pseudomonas aeruginosa</i> conjugative plasmid pUM505 containing virulence and heavy-metal resistance genes. Plasmid. 2011. 66:7-18
R. J. Shea, M. H. Mulks. ohr, Encoding an organic hydroperoxide reductase, is an in vivo-induced gene in <i>Actinobacillus pleuropneumoniae</i> . Infect Immun. 2002. 70:794-802
M. M. Babykin, K. V. Sidoruk, V. V. Zinchenko, L. N. Nefedova, R. Cerff, S. V. Shestakov. [On the involvement of the regulatory gene prqR in the development of resistance to methyl viologen in cyanobacterium Synechocystis sp. PCC6803]. Genetika. 2003. 39:25-32
J. H. Shin, K. Yoshimoto, Y. Ohsumi, J. S. Jeon, G. An. OsATG10b, an autophagosome component, is needed for cell survival against oxidative stresses in rice. Mol Cells. 2009. 27:67-74
Y. S. Kang, Y. Lee, H. Jung, C. O. Jeon, E. L. Madsen, W. Park. Overexpressing antioxidant enzymes enhances naphthalene biodegradation in <i>Pseudomonas</i> sp. strain As1. Microbiology. 2007. 153:3246-54

Fur (ferric uptake regulator) is a prokaryotic transcriptional regulator that controls a large number of genes mainly relate

The ability to induce cellular defense mechanisms in response to environmental challenges is a fundamental property of

Nitric oxide (NO) is emerging as a key regulator of diverse plant cellular processes. A major route for the transfer of NO b

Nitroreductase A catalyzes the divalent reduction of nitro compounds, quinones, and dyes by NADPH. In this paper, nitro

Formation of nonnative disulfide bonds in the cytoplasm, so-called disulfide stress, is an integral component of oxidative

The charged quaternary ammonium compounds--methyl, ethyl and benzyl viologens--generate reactive oxygen species i

A new oxidative stress-responsive bacterial biosensor was constructed using the promoter of the pgi gene fused to the lu

The full-length cDNA of CaAbs1 encodes a presumptive protein of 134 amino acid residues that has homology to a putat

All organisms have defense systems against oxidative stress that include multiple genes of antioxidant defense. These ge

Manganese ions (Mn(2+)) play a crucial role in virulence and protection against oxidative stress in bacterial pathogens. S

Citrus salt-stress associated protein (Cit-SAP) reveals significant sequence homology to mammalian glutathione peroxida

NRAMPs (natural resistance-associated macrophage proteins) have been characterized in mammals as divalent transition

The ability to adapt to acute oxidative stress (e.g. H₂O₂), peroxynitrite, menadione, and paraquat) through transient a

We determined the complete nucleotide sequence of conjugative plasmid pUM505 isolated from a clinical strain of Pseu

Actinobacillus pleuropneumoniae is the causative agent of porcine pleuropneumonia, a disease characterized by pulmon

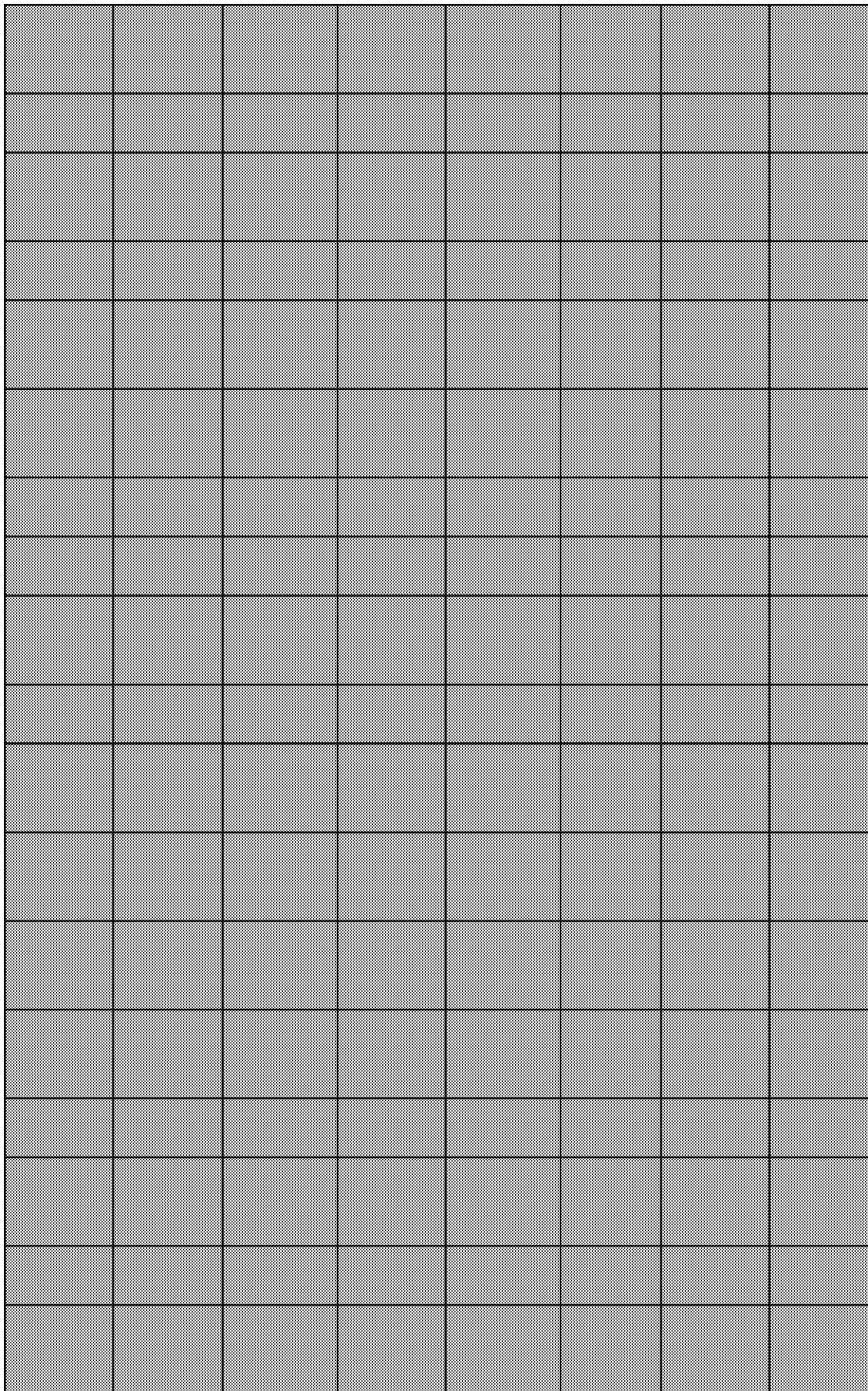
The role of the prqR gene in the regulation of the adaptive response of the cyanobacterium Synechocystis sp. PCC6803 to

Autophagy degrades toxic materials and old organelles, and recycles nutrients in eukaryotic cells. Whereas the studies on

We tested the hypothesis that during metabolism of naphthalene and other substrates by *Pseudomonas* sp. strain As1 ox

Not Relevant

Level 1



2327
2328
2329
2330
2331
2332
2333
2334
2335
2336
2337
2338
2339
2340
2341
2342
2343

H. Y. Yoo, S. S. Kim, H. M. Rho. Overexpression and simple purification of human superoxide dismutase (SOD1) in yeast and its resistance to oxidative stress. <i>J Biotechnol.</i> 1999. 68:29-35
H. S. Yoon, I. A. Lee, H. Lee, B. H. Lee, J. Jo. Overexpression of a eukaryotic glutathione reductase gene from <i>Brassica campestris</i> improved resistance to oxidative stress in <i>Escherichia coli</i> . <i>Biochem Biophys Res Commun.</i> 2005. 326:618-23
A. Szakmary, S. M. Huang, D. T. Chang, P. A. Beachy, M. Sander. Overexpression of a Rrp1 transgene reduces the somatic mutation and recombination frequency induced by oxidative DNA damage in <i>Drosophila melanogaster</i> . <i>Proc Natl Acad Sci U S A.</i> 1996. 93:1607-12
J. Moskovitz, E. Flescher, B. S. Berlett, J. Azare, J. M. Poston, E. R. Stadtman. Overexpression of peptide-methionine sulfoxide reductase in <i>Saccharomyces cerevisiae</i> and human T cells provides them with high resistance to oxidative stress. <i>Proc Natl Acad Sci U S A.</i> 1998. 95:14071-5
Y. H. Kim, M. H. Yu. Overexpression of reactive cysteine-containing 2-nitrobenzoate nitroreductase (NbaA) and its mutants alters the sensitivity of <i>Escherichia coli</i> to reactive oxygen species by reprogramming a regulatory network of disulfide-bonded proteins. <i>J Proteome Res.</i> 2012. 11:3219-30
T. Fujibe, H. Saji, M. K. Watahiki, K. T. Yamamoto. Overexpression of the RADICAL-INDUCED CELL DEATH1 (RCD1) gene of <i>Arabidopsis</i> causes weak rcd1 phenotype with compromised oxidative-stress responses. <i>Biosci Biotechnol Biochem.</i> 2006. 70:1827-31
K. Hakkila, T. Antal, A. U. Rehman, J. Kurkela, H. Wada, I. Vass, E. Tyystjarvi, T. Tyystjarvi. Oxidative stress and photoinhibition can be separated in the cyanobacterium <i>Synechocystis</i> sp. PCC 6803. <i>Biochim Biophys Acta.</i> 2014. 1837:217-25
S. Belkin, D. R. Smulski, A. C. Vollmer, T. K. Van Dyk, R. A. LaRossa. Oxidative stress detection with <i>Escherichia coli</i> harboring a katG::lux fusion. <i>Appl Environ Microbiol.</i> 1996. 62:2252-6
N. Mehterov, S. Balazadeh, J. Hille, V. Toneva, B. Mueller-Roeber, T. Gechev. Oxidative stress provokes distinct transcriptional responses in the stress-tolerant atr7 and stress-sensitive loh2 <i>Arabidopsis thaliana</i> mutants as revealed by multi-parallel quantitative real-time PCR analysis of ROS marker and antioxidant genes. <i>Plant Physiol Biochem.</i> 2012. 59:20-9
E. R. Rocha, T. Selby, J. P. Coleman, C. J. Smith. Oxidative stress response in an anaerobe, <i>Bacteroides fragilis</i> : a role for catalase in protection against hydrogen peroxide. <i>J Bacteriol.</i> 1996. 178:6895-903
M. E. Tosello, M. S. Biasoli, A. G. Luque, H. M. Magaro, A. R. Krapp. Oxidative stress response involving induction of protective enzymes in <i>Candida dubliniensis</i> . <i>Med Mycol.</i> 2007. 45:535-40
S. Ito-Kuwa, K. Nakamura, S. Aoki, T. Osafune, V. Vidotto, K. Pienthaweechai. Oxidative stress sensitivity and superoxide dismutase of a wild-type parent strain and a respiratory mutant of <i>Candida albicans</i> . <i>Med Mycol.</i> 1999. 37:307-14
N. S. Jakubovics, A. W. Smith, H. F. Jenkinson. Oxidative stress tolerance is manganese (Mn(2+)) regulated in <i>Streptococcus gordonii</i> . <i>Microbiology.</i> 2002. 148:3255-63
W. F. Li, J. Ji, G. Wang, H. Y. Wang, B. L. Niu, T. L. Josine. Oxidative stress-resistance assay for screening yeast strains overproducing heterologous proteins. <i>Genetika.</i> 2011. 47:1175-83
S. Mendez-Alvarez, K. Rufenacht, R. I. Eggen. The oxidative stress-sensitive yap1 null strain of <i>Saccharomyces cerevisiae</i> becomes resistant due to increased carotenoid levels upon the introduction of the <i>Chlamydomonas reinhardtii</i> cDNA, coding for the 60S ribosomal protein L10a. <i>Biochem Biophys Res Commun.</i> 2000. 267:953-9
J. H. Lee, C. H. Youn, B. C. Kim, M. B. Gu. An oxidative stress-specific bacterial cell array chip for toxicity analysis. <i>Biosens Bioelectron.</i> 2007. 22:2223-9
A. A. Olczak, J. W. Olson, R. J. Maier. Oxidative-stress resistance mutants of <i>Helicobacter pylori</i> . <i>J Bacteriol.</i> 2002. 184:3186-93

The structural gene of human Cu/Zn superoxide dismutase (hSOD1) was cloned into a yeast expression vector containing

Glutathione reductase (GR) plays an essential role in a cell's defense against reactive oxygen metabolites by sustaining th

Recombination repair protein 1 (Rrp1) includes a C-terminal region homologous to several DNA repair proteins, including

The yeast peptide-methionine sulfoxide reductase (MsrA) was overexpressed in a *Saccharomyces cerevisiae* null mutant

The effects of redox-sensitive proteins on *Escherichia coli* were investigated by overexpressing *Pseudomonas* 2-nitroben

rcd1 is a mutant of *Arabidopsis thaliana* that is more resistant to methyl viologen, but more sensitive to ozone than the w

Roles of oxidative stress and photoinhibition in high light acclimation were studied using a regulatory mutant of the cyan

A plasmid containing a transcriptional fusion of the *Escherichia coli* katG promoter to a truncated *Vibrio fischeri lux* oper

The *Arabidopsis thaliana* *atr7* mutant is tolerant to oxidative stress induced by paraquat (PQ) or the catalase inhibitor am

Survival of *Bacteroides fragilis* in the presence of oxygen was dependent on the ability of bacteria to synthesize new prot

Candida dubliniensis is a yeast species closely related to *Candida albicans*, but in contrast to *C. albicans*, limited informat

It is important to know responses of the pathogenic fungi to reactive oxygen species by which hosts protect themselves a

The Sca permease in the oral bacterium *Streptococcus gordonii* is a member of a family of ATP-binding cassette (ABC)-ty

Many natural proteins have been developed into drugs and produced for direct application. Identifying improved hosts t

The *Saccharomyces cerevisiae* *yap1* null strain was transformed with a *Chlamydomonas reinhardtii* cDNA expression libra

An oxidative stress-specific bacterial cell array chip was fabricated and implemented in the analysis of various different c

Within a large family of peroxidases, one member that catalyzes the reduction of organic peroxides to alcohols is known

Not Relevant

Not Relevant